

APPROVED	C.B. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

CCACC ATG GCT CTG CAG ATC CCC AGC CTC CTC TCA GCT GCT GTG GTG CTG ATG GTG CTG AGC AGC CCA AGG  
M A L Q I P S L L L L S A A V V V L M V L S S P R  
KOZAK \_\_\_\_\_ IA<sup>d</sup> β CHAIN SIGNAL PEPTIDE \_\_\_\_\_  
CONSENSUS

ACC TTA AGT ATC TCT CAG GCT GTT CAC GCT GCT CAC GCT GAA ATC AAC GAA GCT GGT CGT  
T L S ↑ I S Q A V H A A H A E I N E A G R  
\_\_\_\_\_ OVA 323-339 PEPTIDE \_\_\_\_\_

GCT AGC GGA GGG GGC GGA AGC GGC GGA GGG GGA AAC TCC GAA AGG // AGC CCC ATC ACT GTG GAG TGG  
A S G G G G S G G G G G N S E R // S P I T V E W  
\_\_\_\_\_ PEPTIDE LINKER \_\_\_\_\_ aa1 \_\_\_\_\_ IA<sup>d</sup> β1-β2 DOMAINS \_\_\_\_\_ aa189

ACT AGT GGT GGC GGT GGC AGC GGC GGT GGT GGT TCC GGT GGC GGC GGT TCT GGC GGT GGC GGT TCC TCG AGT  
T S G G G G S G G G G G S G G G S G G G S S S  
\_\_\_\_\_ SINGLE CHAIN LINKER \_\_\_\_\_

GAA GAC GAC ATT // CCA GGG CCT TTA TGA  
E D D I // P G P L • STOP  
aa1 \_\_\_\_\_ IA<sup>d</sup> α CHAIN \_\_\_\_\_

FIG. 1

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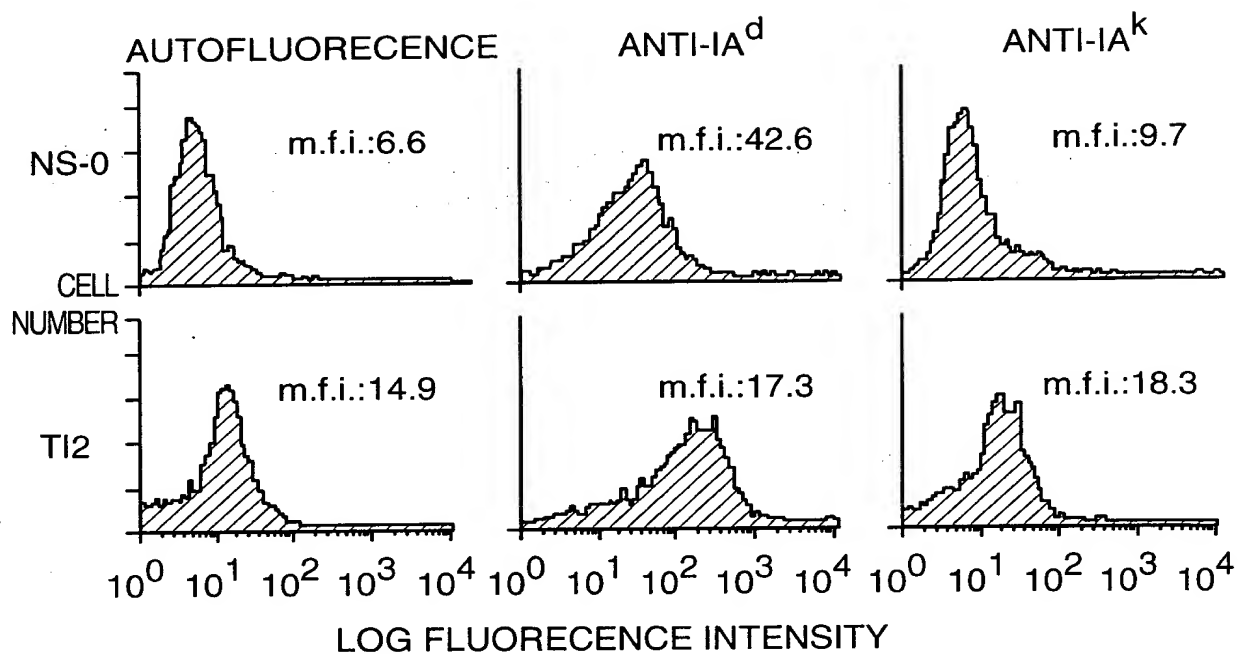


FIG. 2A

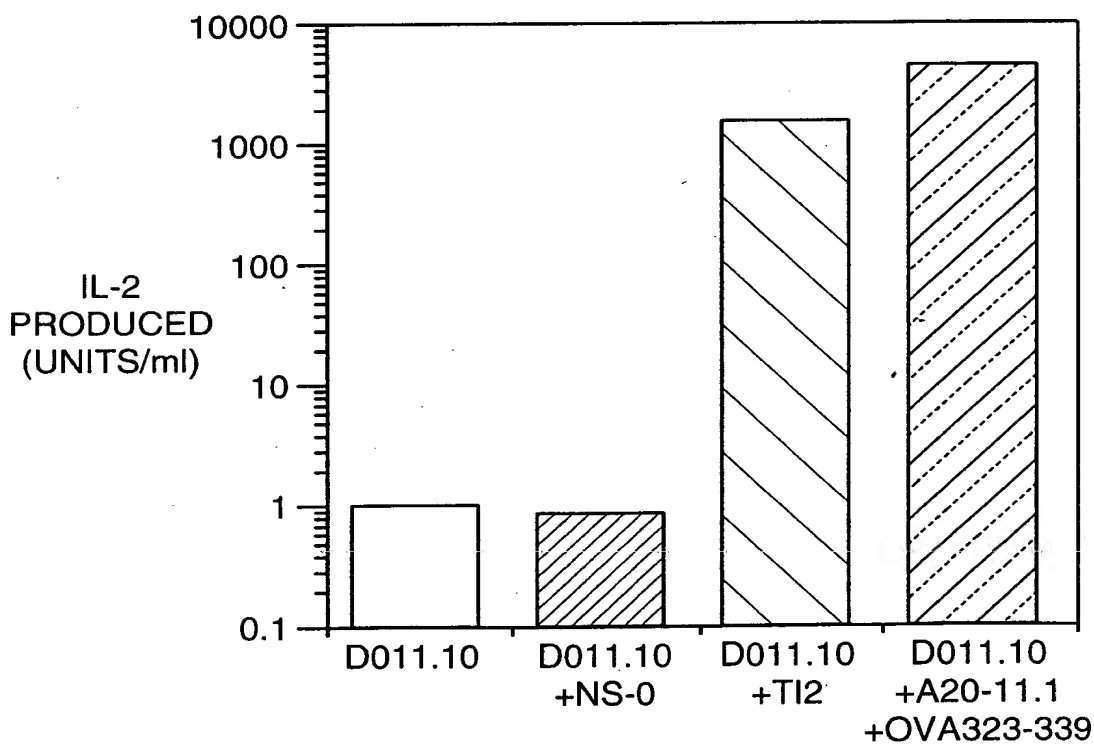


FIG. 2B

FOR FIG. 2A

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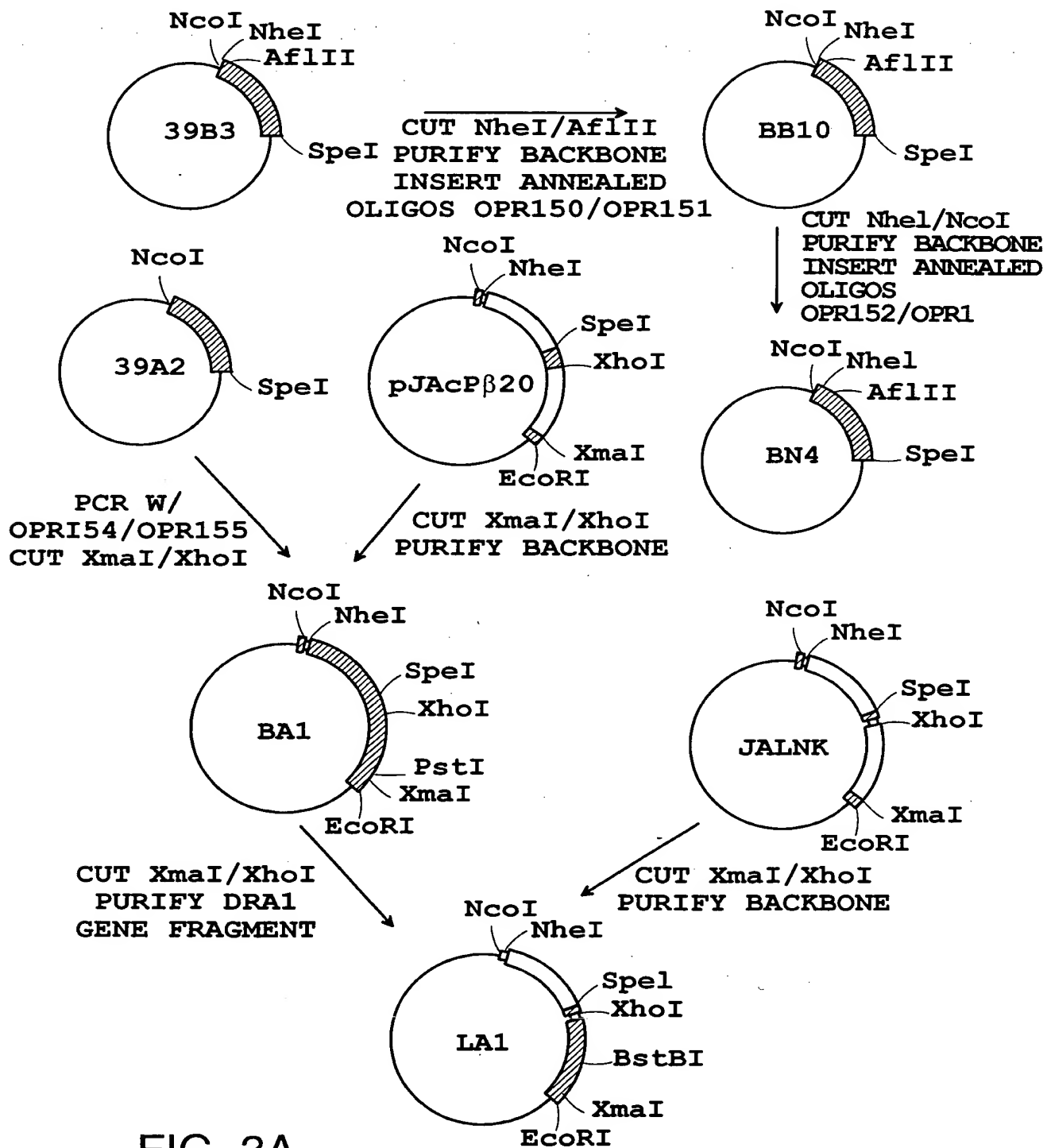


FIG. 3A

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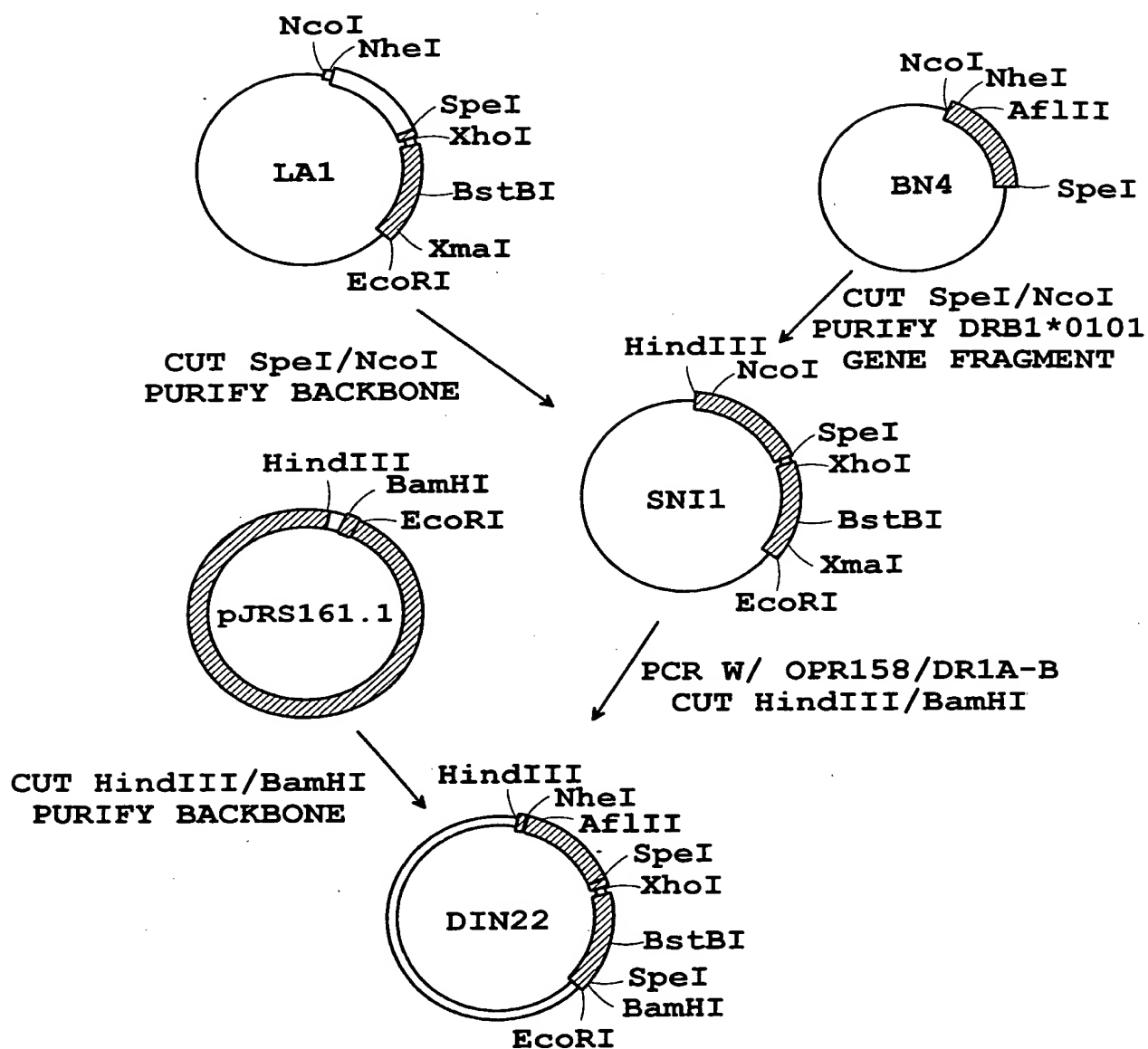


FIG. 3B

FIG. 3B

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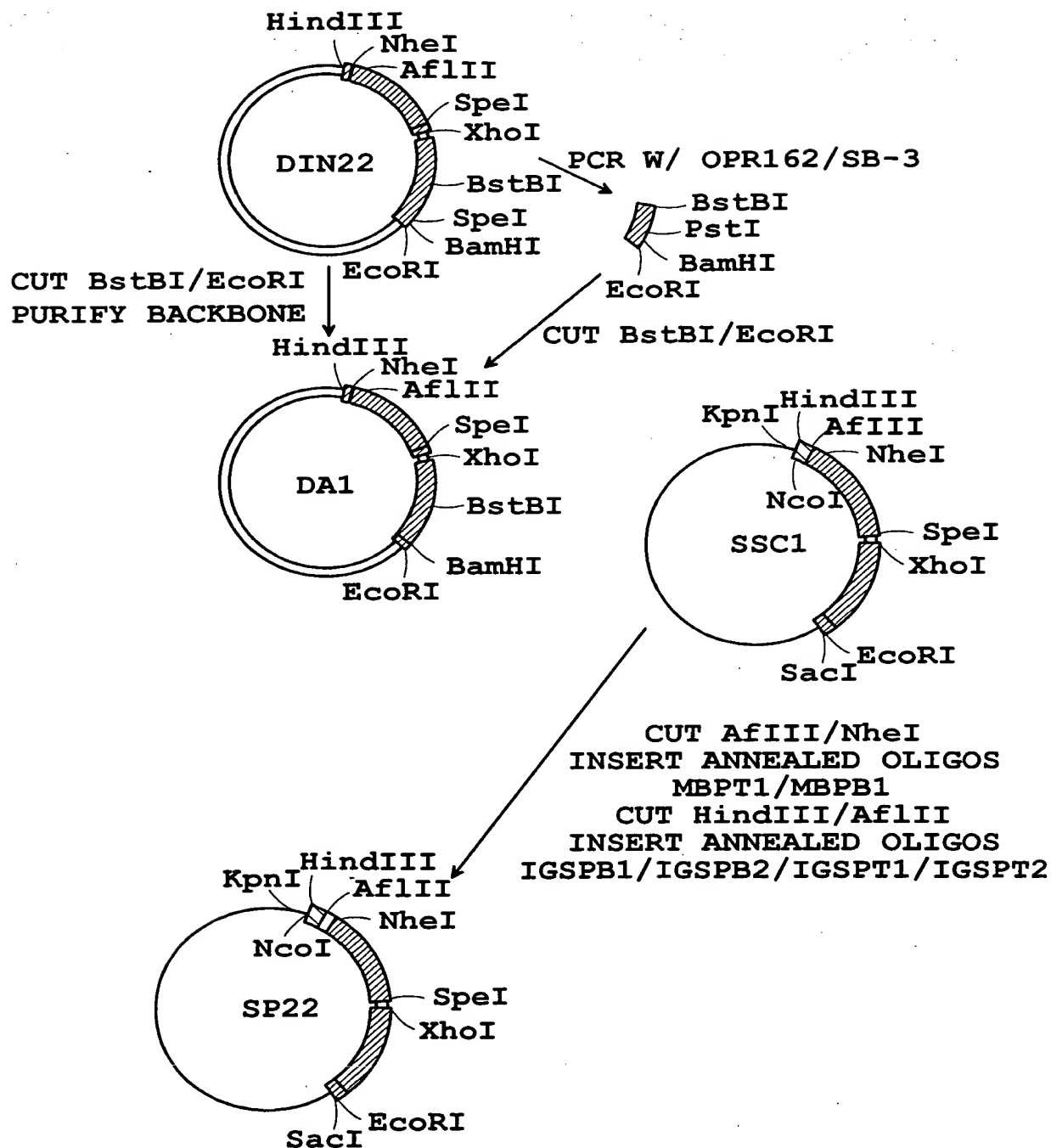


FIG. 3C

TOP SECRET

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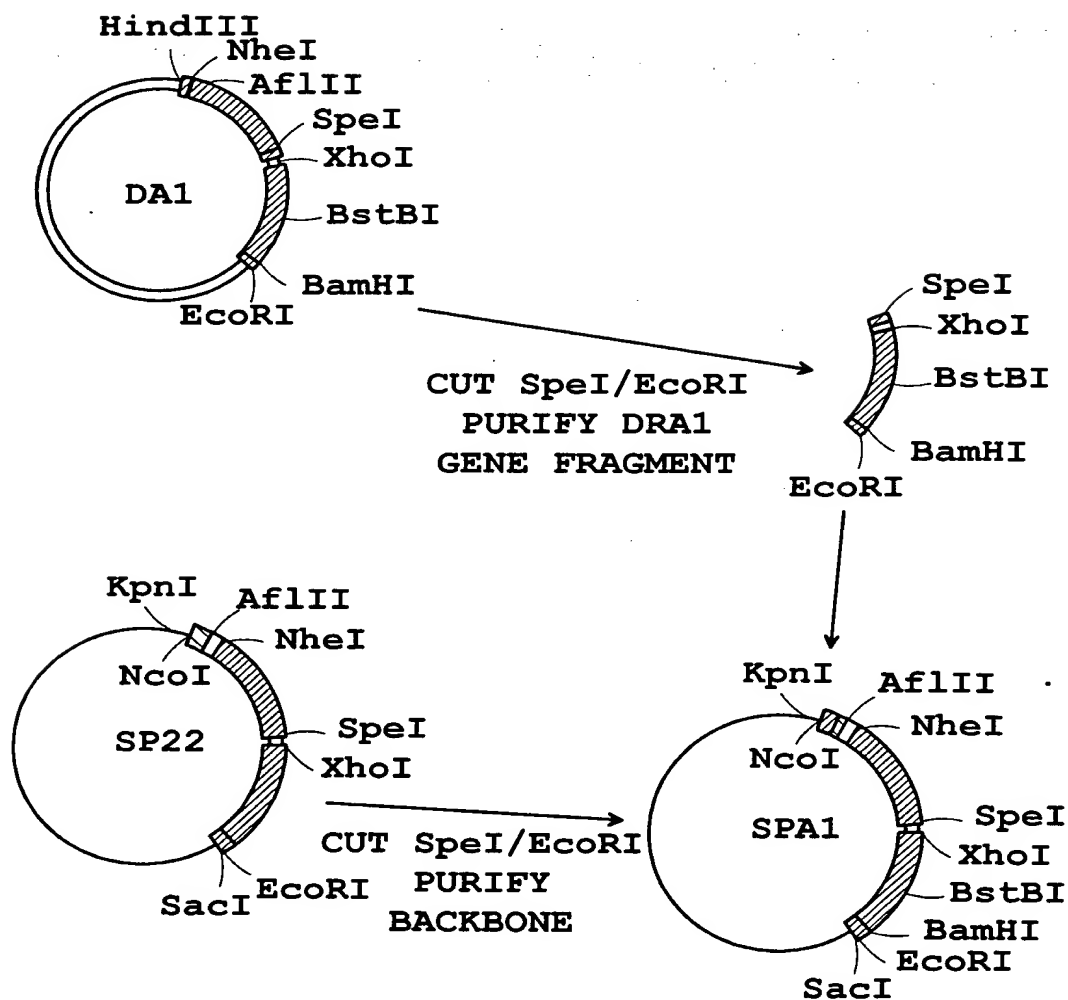


FIG. 3D

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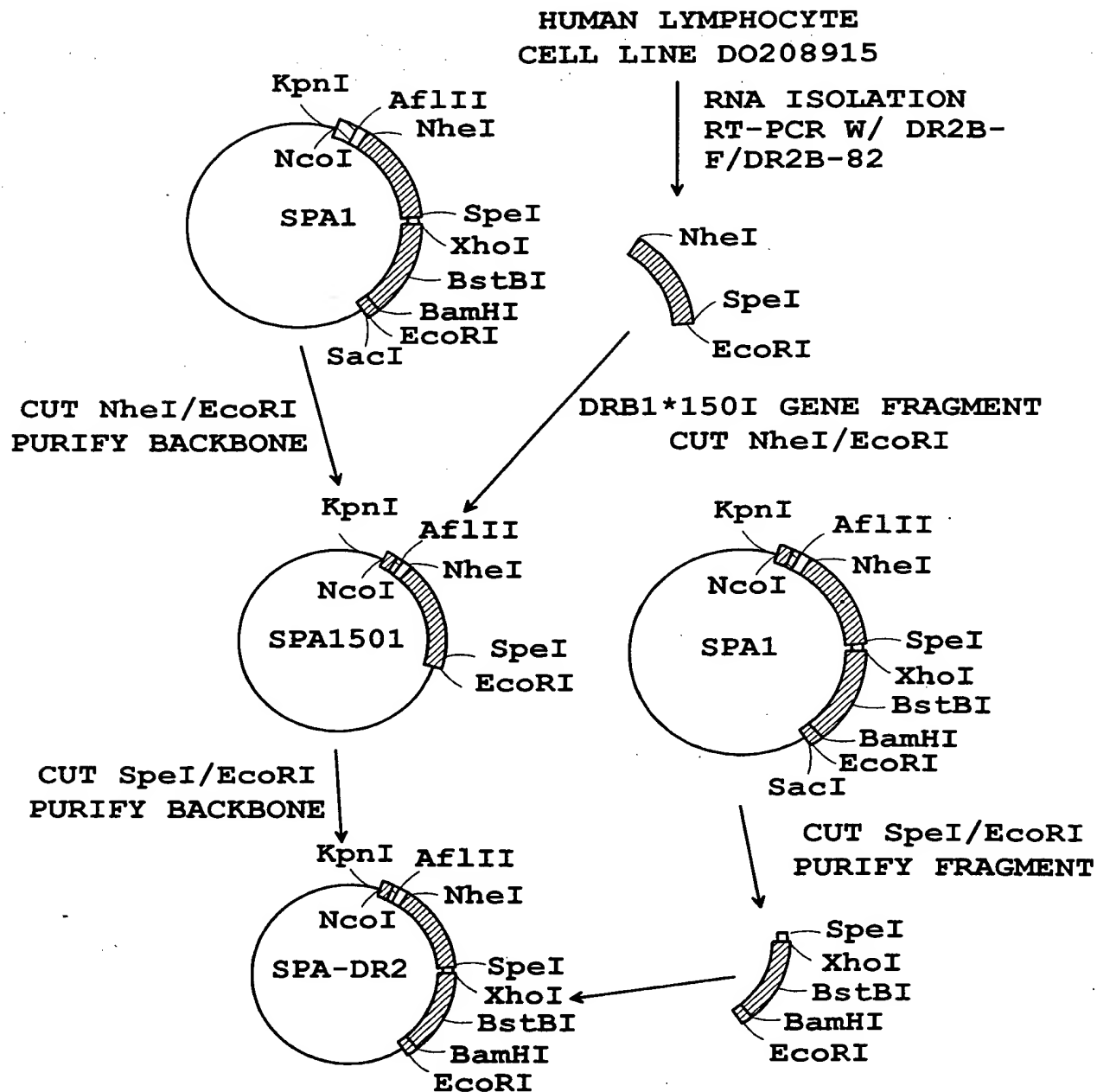


FIG. 3E

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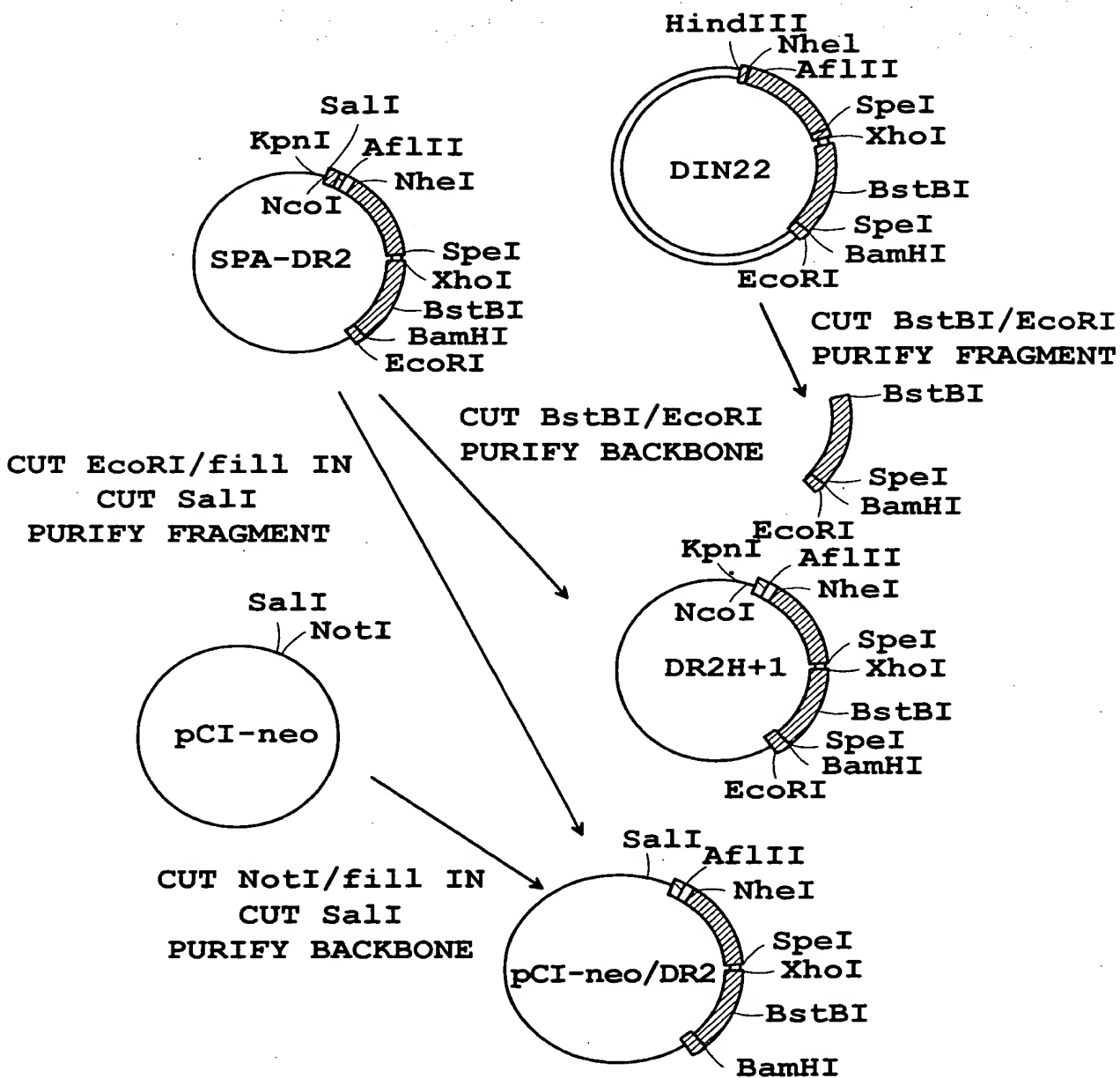


FIG. 3F



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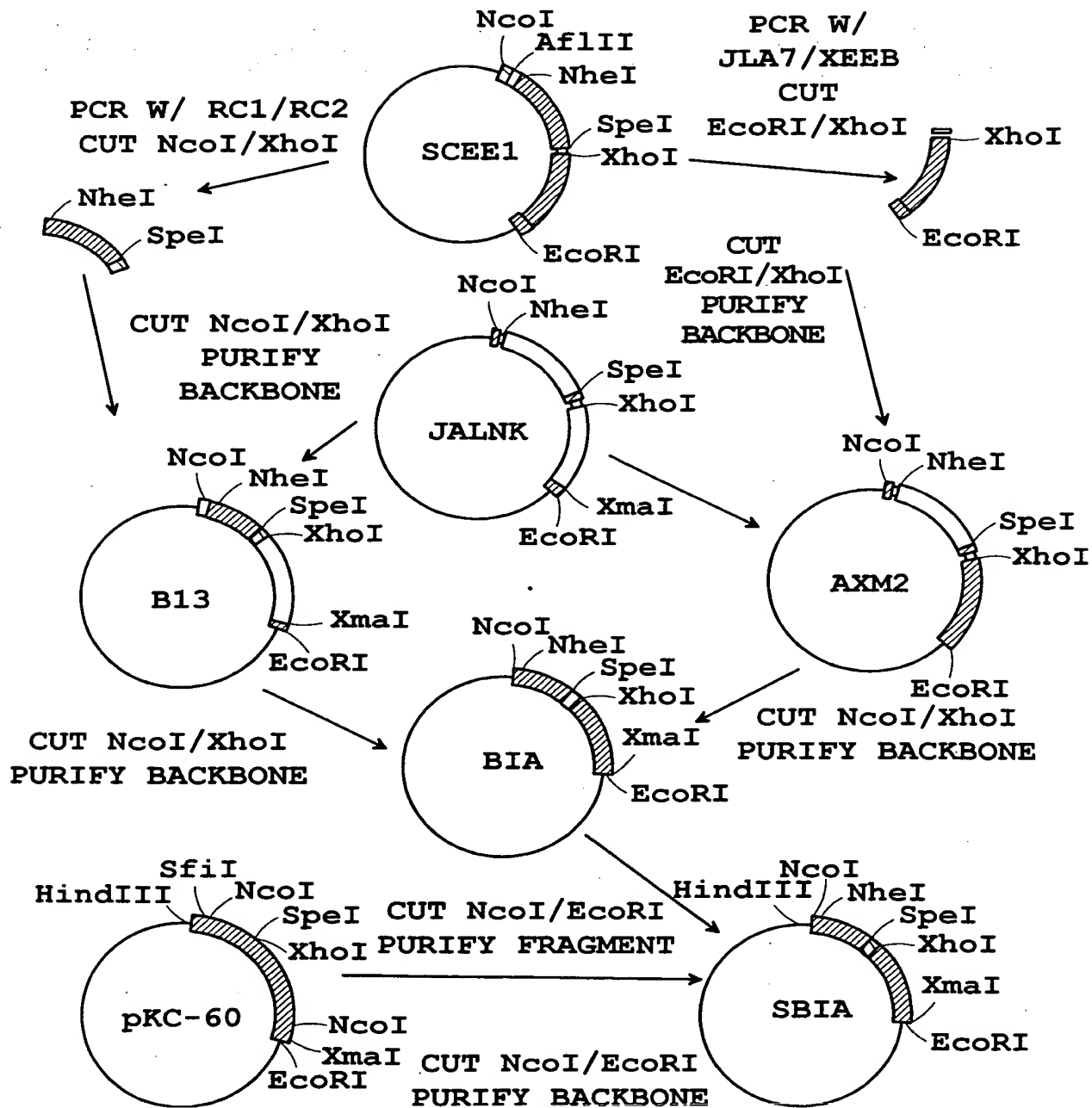


FIG. 3G

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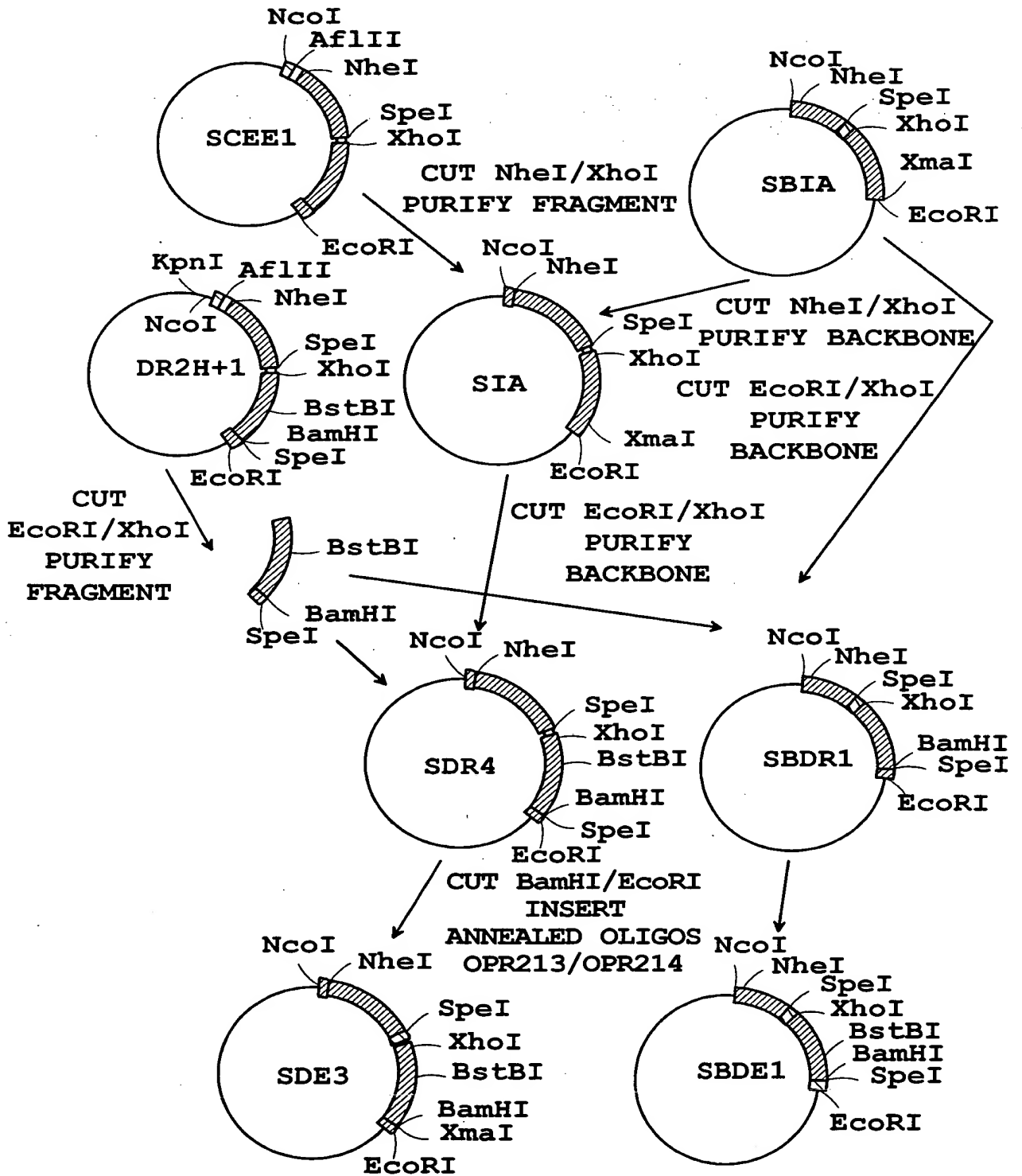


FIG. 3H

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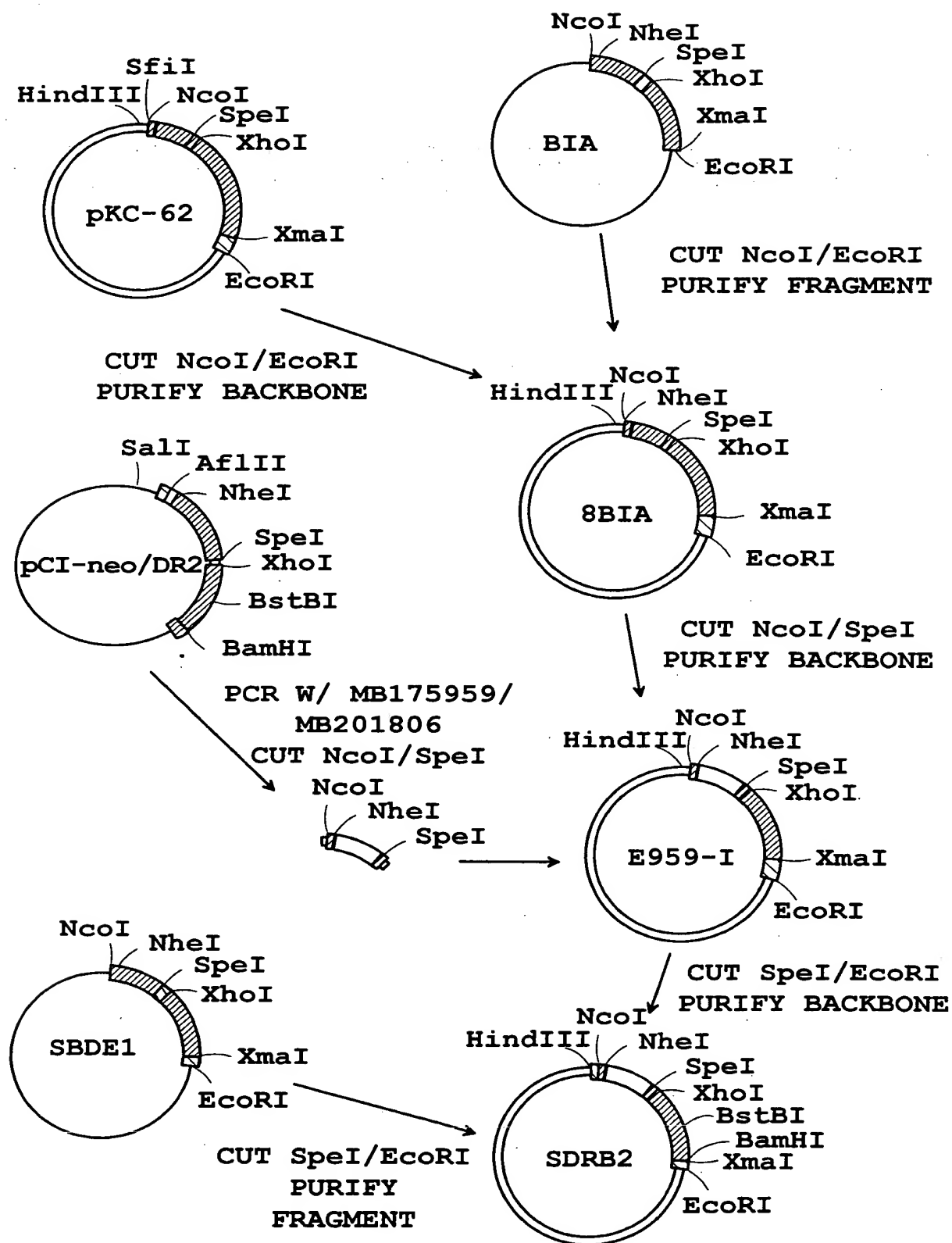


FIG. 3I

FIG. 3I

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FIG. 3J

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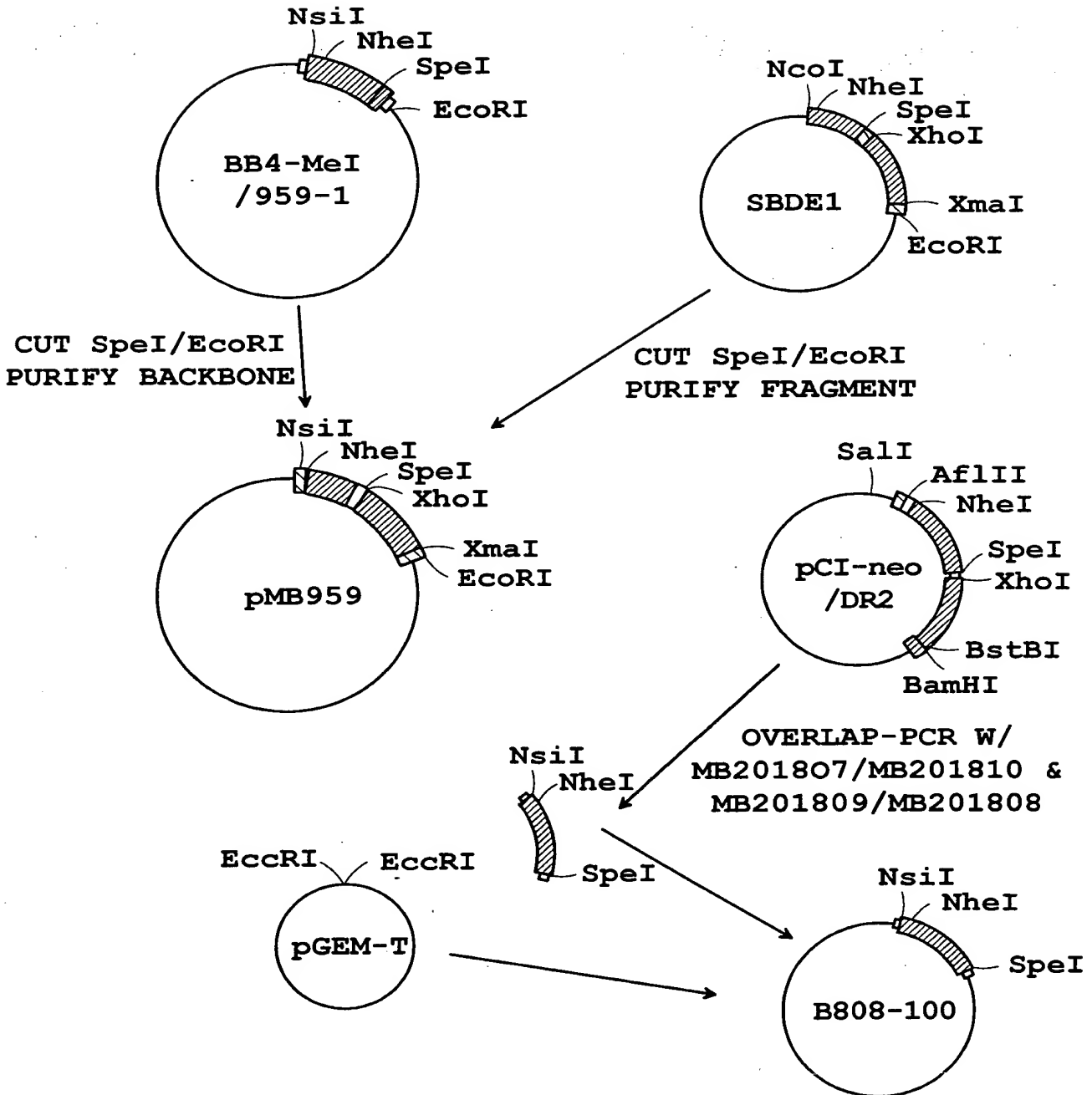


FIG. 3K

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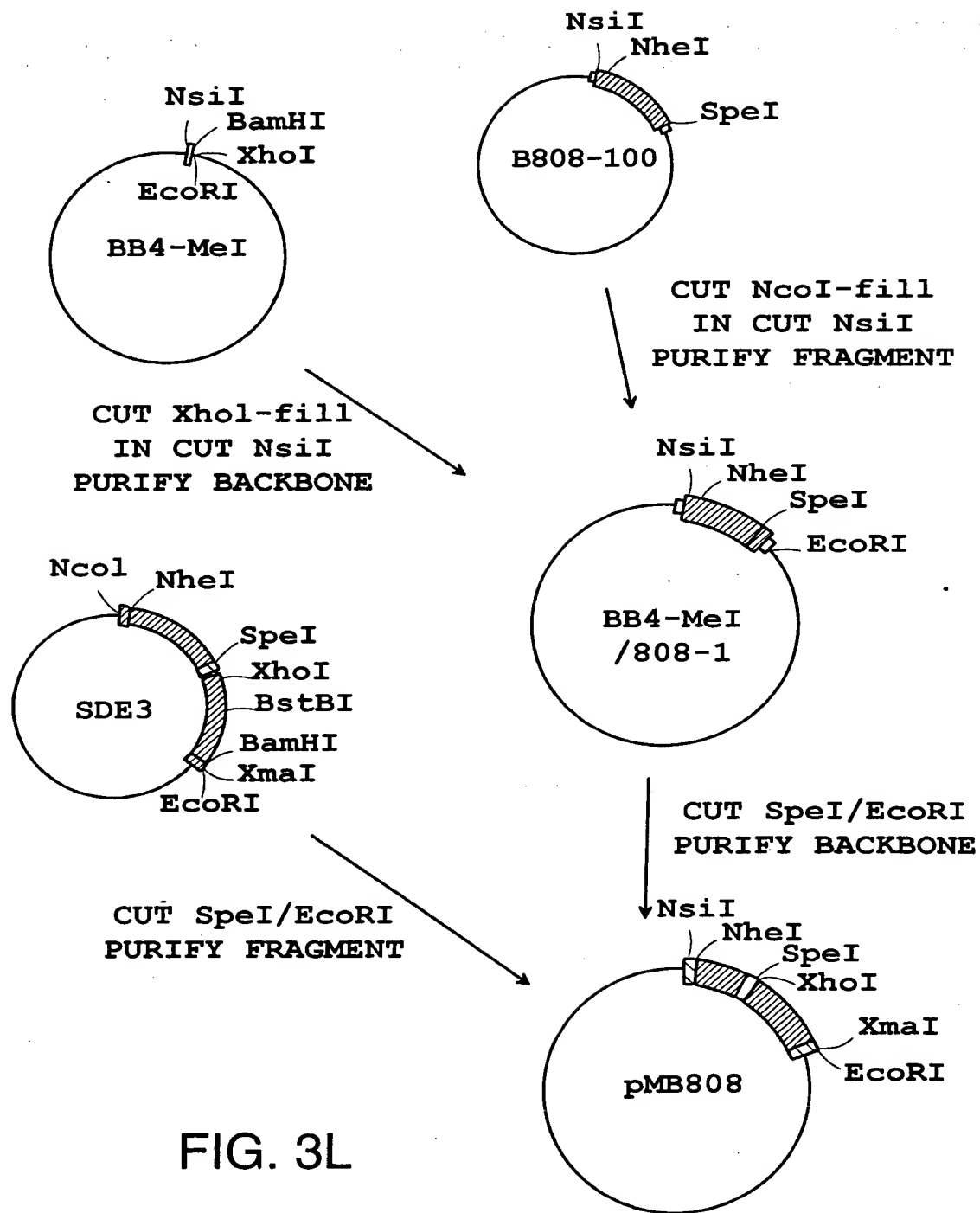


FIG. 3L

TOP SECRET

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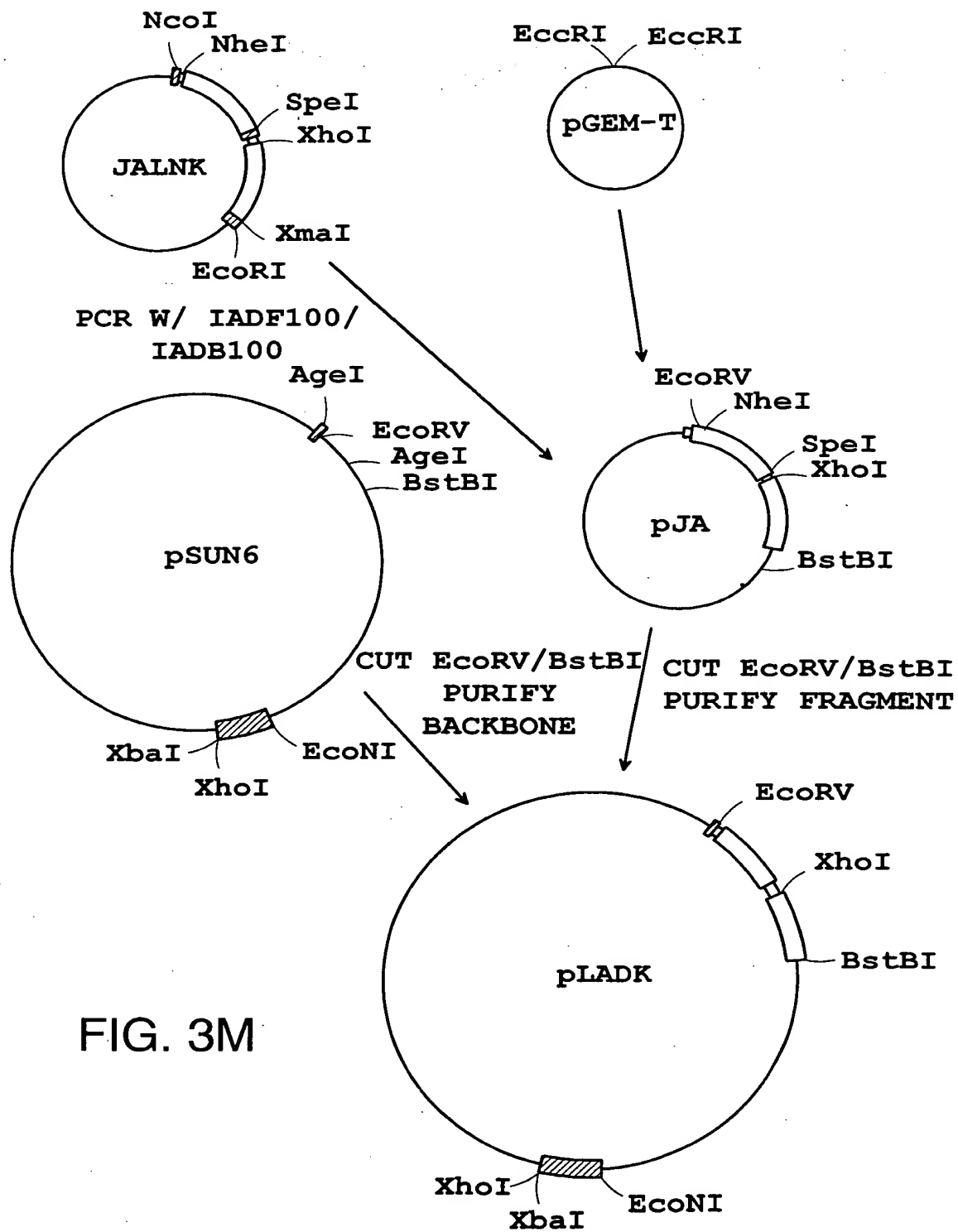


FIG. 3M

00766338-01100  
FIG. 3M

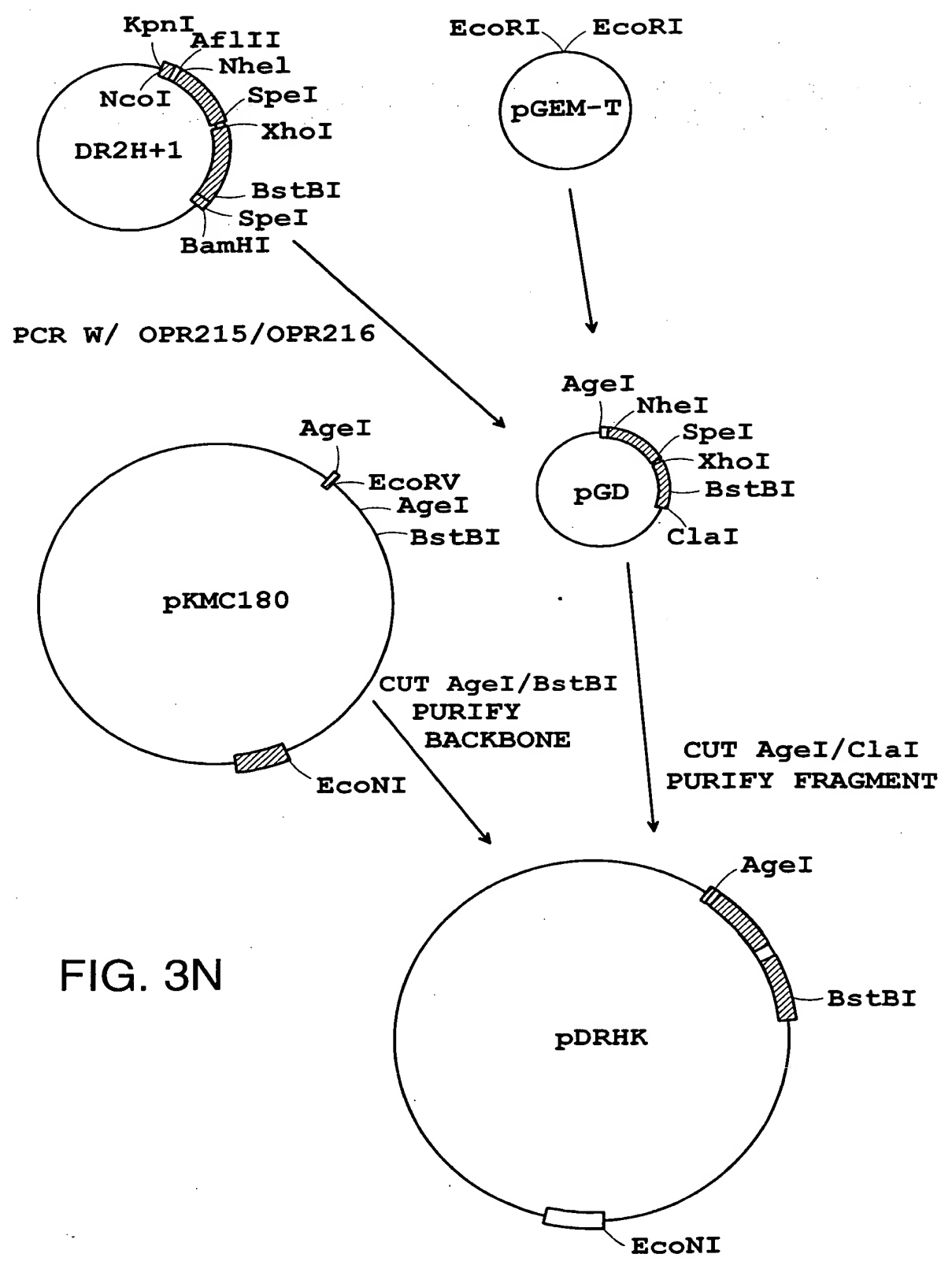


FIG. 3N

FIG. 3N



sc-1A<sup>d</sup>/PEPTIDE FUSION

SP	PEP	L1	IA <sup>d</sup> β1-β2	L2	IA <sup>d</sup> α1-α2
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sc-1A<sup>d</sup>/PEPTIDE-TAG FUSION

SP	PEP	L1	IA <sup>d</sup> β1-β2	L2	IA <sup>d</sup> α1-α2	EE
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FIG. 4A

sc-1A<sup>d</sup>TM/PEPTIDE FUSION

SP	PEP	L1	IA <sup>d</sup> β1-β2	L2	IA <sup>d</sup> α1-α2	IA <sup>d</sup> αTM-Oγ
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sc-1A<sup>d</sup>/PEPTIDE-C<sub>L</sub> FUSION

SP	PEP	L1	IA <sup>d</sup> β1-β2	L2	IA <sup>d</sup> α1-α2	IgG C <sub>L</sub>
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sc-DR2/PEPTIDE-TAG FUSION

SP	PEP	L1	DRB1*1501 β1-β2	L2	DRAI <sub>α</sub> 1-α2	EE
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sc-DR2-β2/PEPTIDE FUSION

SP	PEP	L1	DRB1*1501 β1	L2	DRAI <sub>α</sub> 1-α2	EE
----	-----	----	--------------	----	------------------------	----

FIG. 4B

sc-DR2 MOD β2/PEPTIDE FUSION

SP	PEP	L1	DRB1*1501 β1 MOD β2	L2	DRAI <sub>α</sub> 1-α2	EE
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sc-DR2/PEPTIDE-C<sub>L</sub> FUSION

SP	PEP	L1	DRB1*1501 β1-β2	L2	DRAI <sub>α</sub> 1-α2	IgG C <sub>L</sub>
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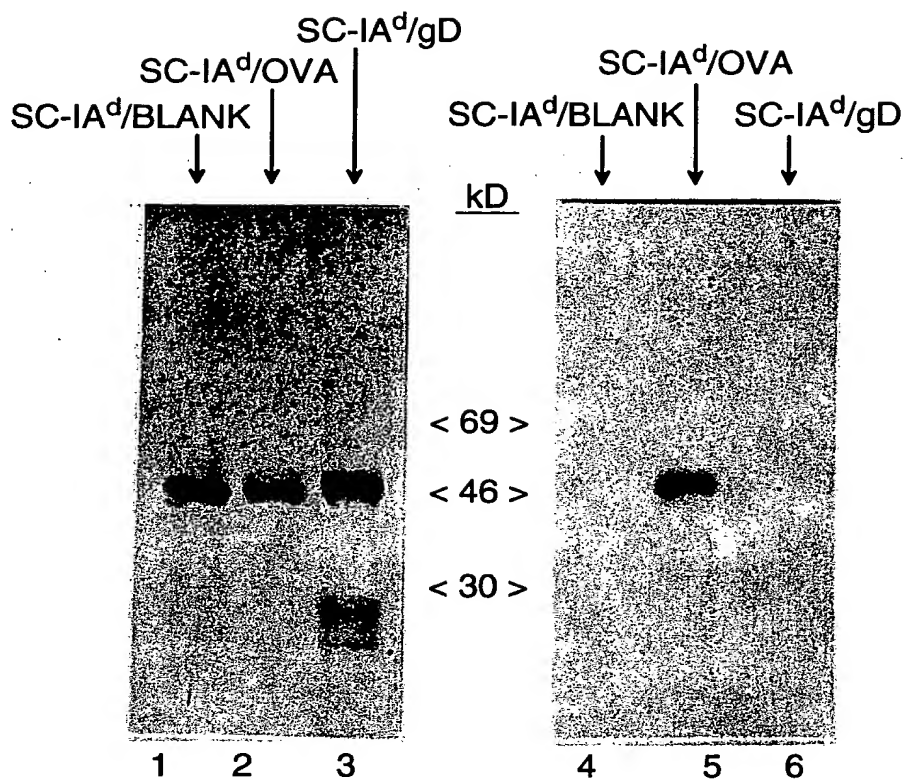


FIG. 5A

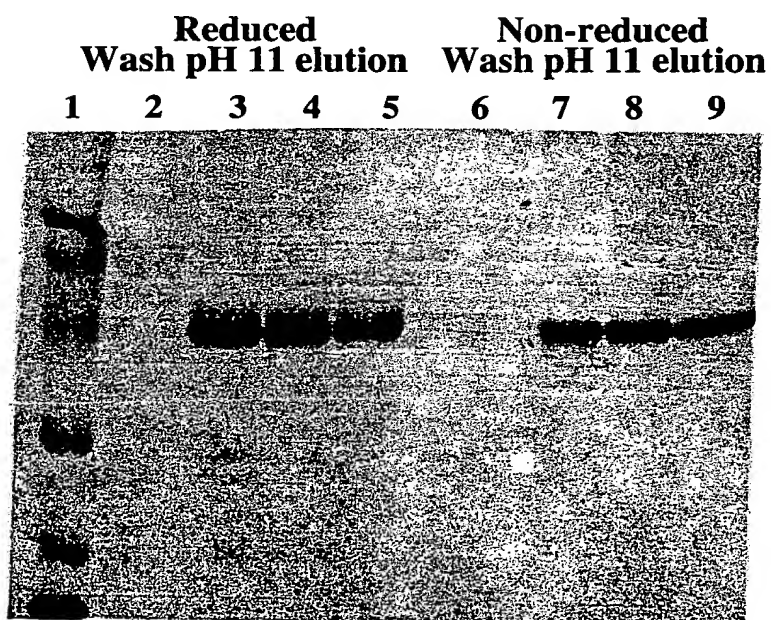
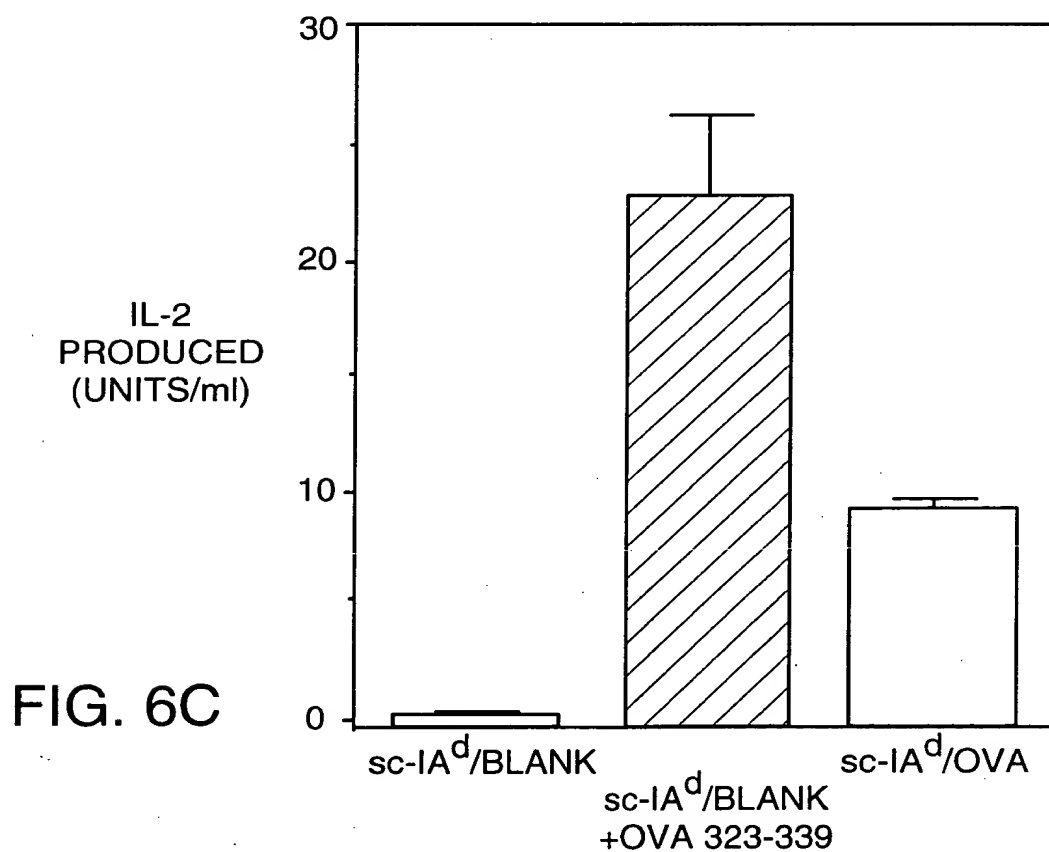
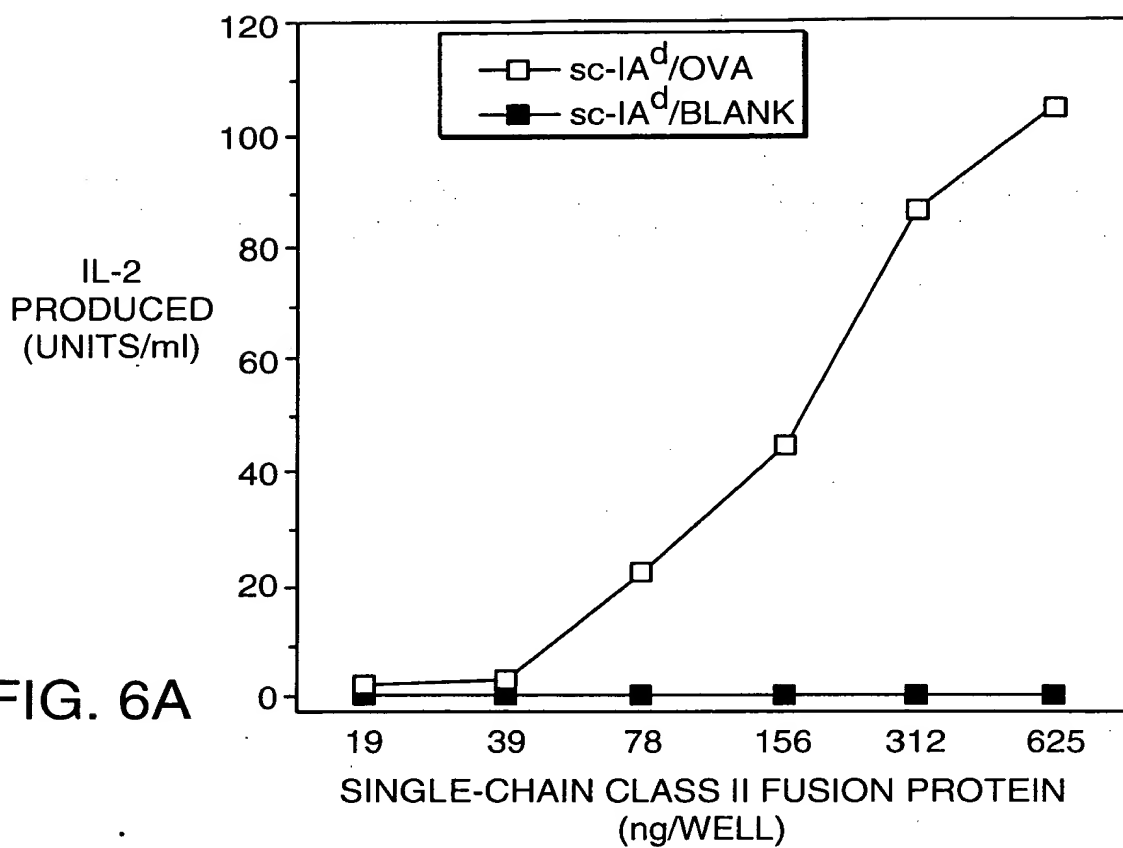


FIG. 5B

FIG. 5A

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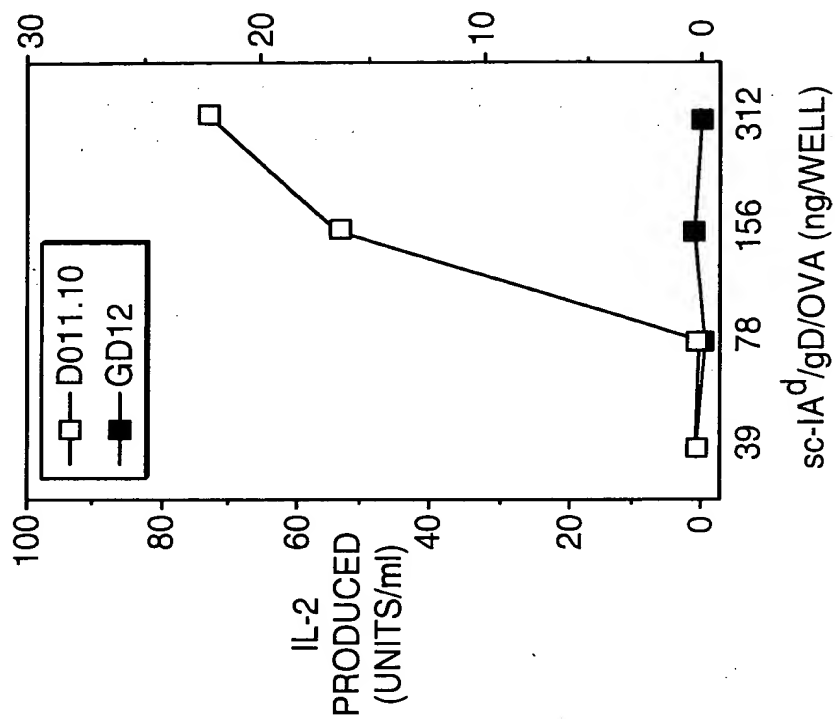


FIG. 6B-2

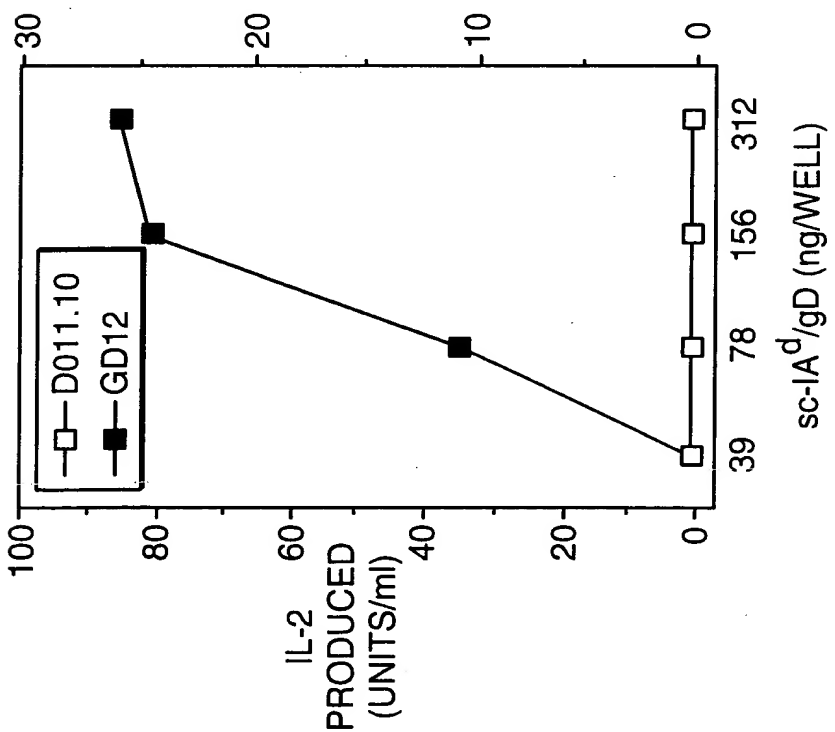
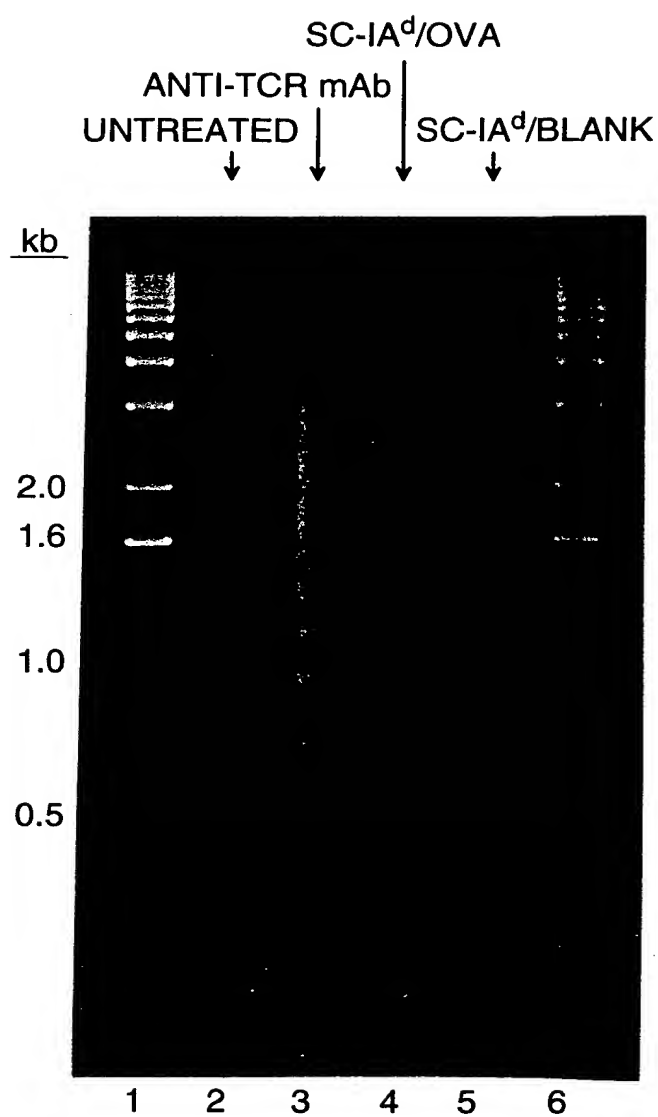


FIG. 6B-1



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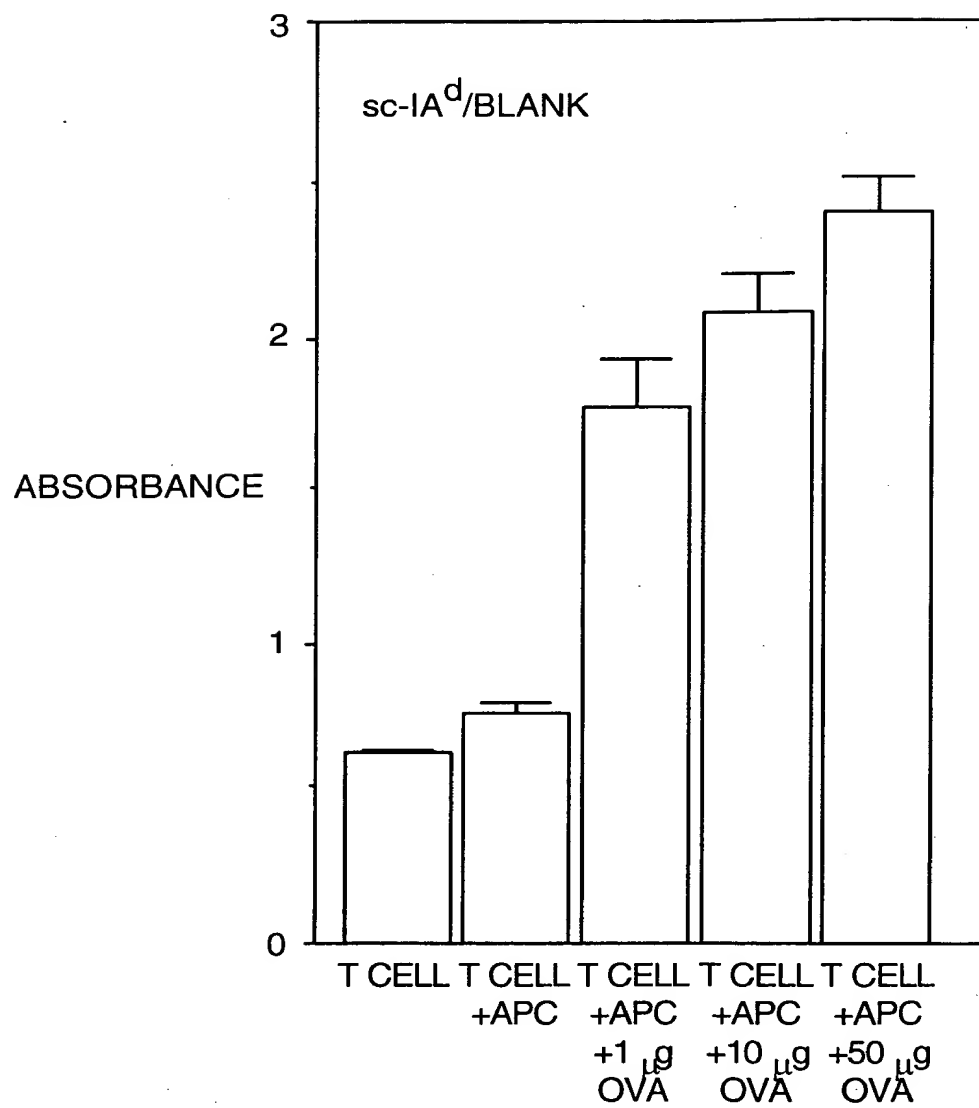


FIG. 8A

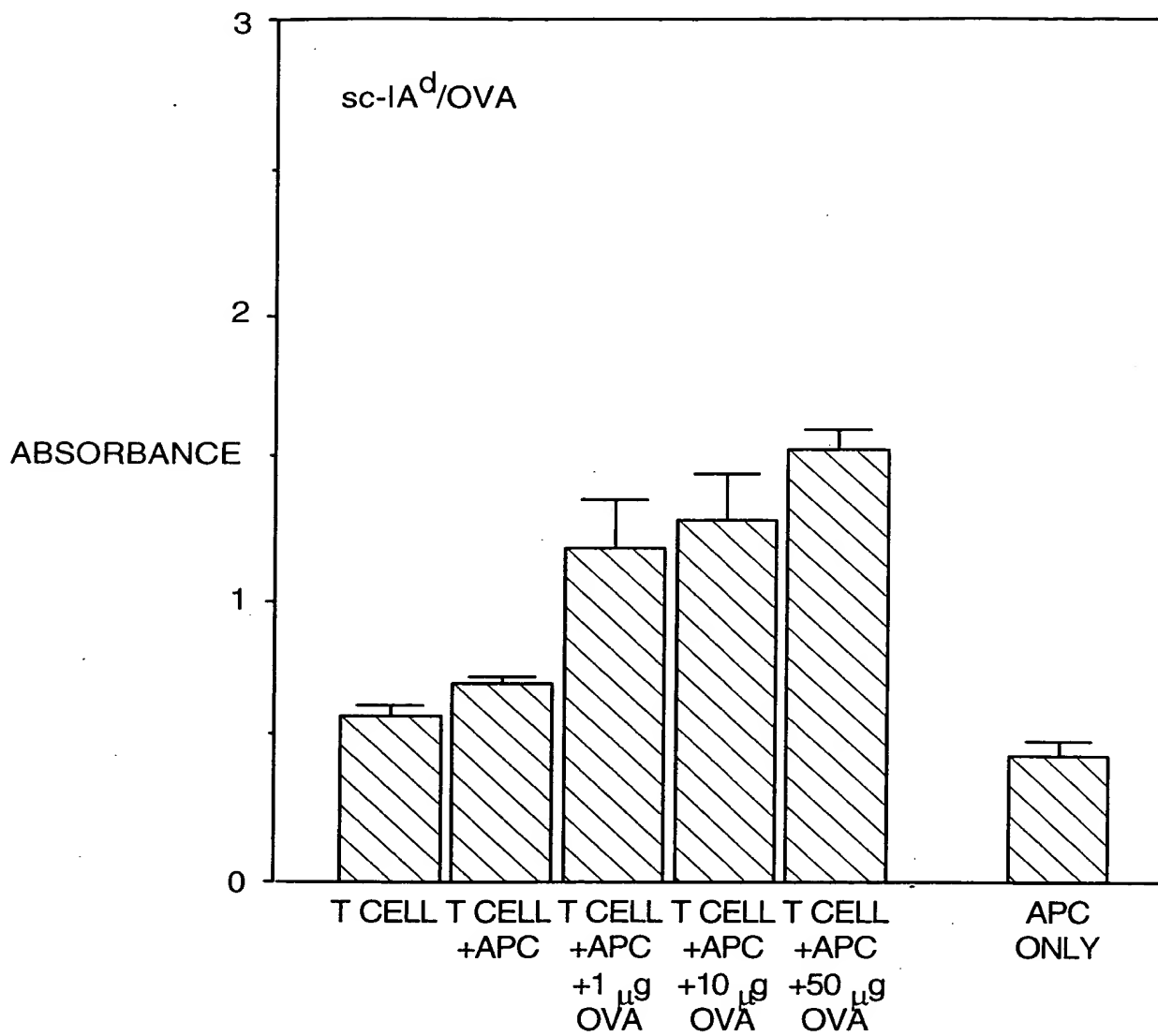


FIG. 8B

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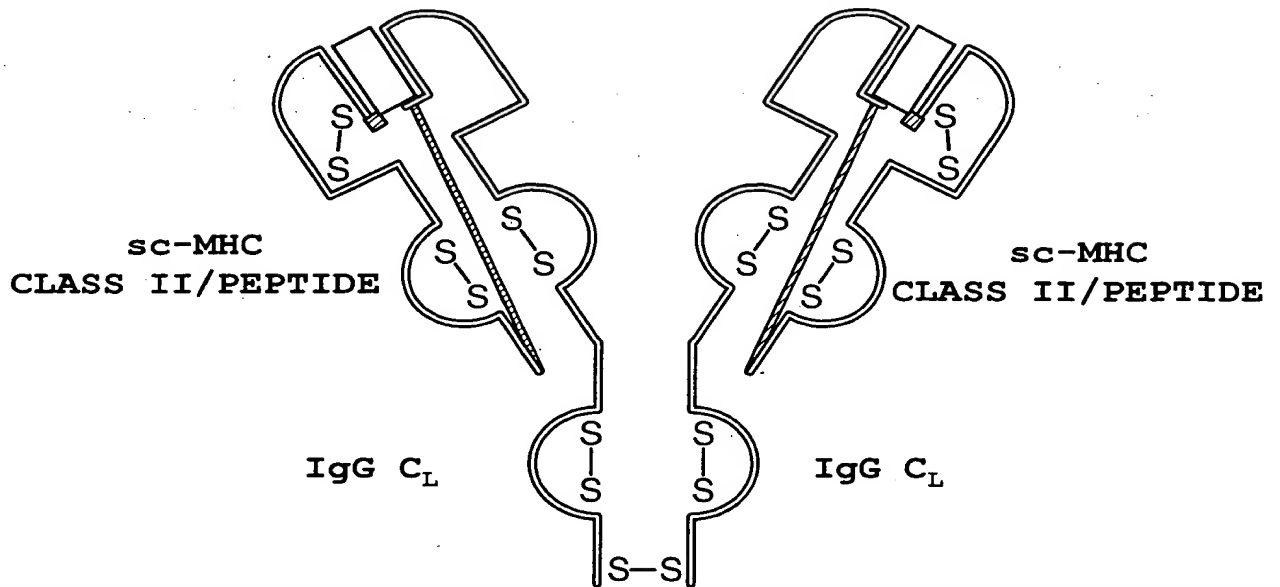


FIG. 9A

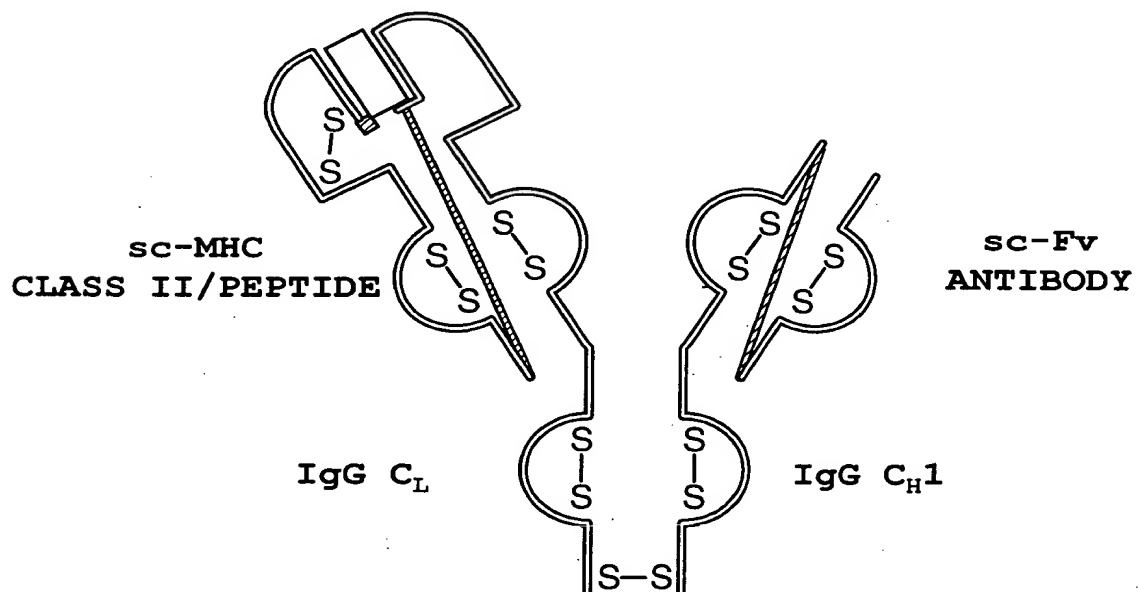


FIG. 9B

FIG. 9A



APPROVED	O.C. FIG.	
BY	DATE	SUBCLASS
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SEQ No.

CCACCATG	1
OPR132 5'-CCCCCAAGCTTCCCGGGCCACCATGGCTCTGCAGATCCCCAGC-3'	2
OPR133 5'-CCCCCACTTAAGGTCCTTGGGCTGCTCAGCACC-3'	3
OPR102 5'-GGGGGGGCCATGGCCGGAACTCCGAAAGGCATTTTCG-3'	4
OPR104 5'-GCGGCGACTAGTCCACTCCACAGTGATGGGGC-3'	5
OPR100 5'-GGGGGGGCCATGGCCGAAGACGACATTGAGGCCGAC-3'	6
OPR101 5'-GCGGCGACTAGTCCAGTGTTTCAGAACCGGCTC-3'.	7
IADF100 5'-GGGGGGGATATCTCTCAGGCTGTTACGCTG-3'	8
IADB100 5'-GGGGGGGTTTCGAAAAGTGTA CTACGGGGGGCTGGAATCTCAGGTTC-3'	9
OPR145 5'-GGGGGGGCTCGAGTATCAAAGAAGAACATGTGATCATC-3'	10
DR1A-B 5'-GCGGCGGGATCCGTTCTCTGTAGTCTCTGGGAGAGG-3'	11
OPR203000 5'-GATCCGAGGAAGAAGAGTACATGCCCATGGAACCCGGGTGAG-3'	12
OPR203001 5'-AATTCTCACCCGGGTTCCATCGGCATGTACTCTTCTTCCTCG-3'	13
DR2B-F 5'-CCCCCGCTAGCGGAGGGGGCGGAAGCGGCGGAGGGGGGGGACA CCCGACCACGTTTCCTGTGGCAGCCTAAGAGG-3'	14
DR2B-B2 5'-CCCCCGAATTCCCCACTAGTCCATTCCACTGTGAGAGGGCTTGTC AC-3'	15
MB201806 5'-GGGGGGGGCCATGGCCTACGACGAGAACCCCGTGGTG-3'	16
MB175959 5'-GGGGGGGACTAGTTCGCCGCTGCACTGTGAAGC-3'	17
MB201807 5'-GGGGGGGTATGCATACGACGAGAACCCCGTGGTG-3'	18
MB201808 5'-GGGGGGGACTAGTTCCACTTCGAGGAACTGTTTCC-3'	19
MB201809 5'-CCTCCTGGTCTCCTCTGTGAGTGG-3'	20
MB201810 5'-CCACTCACAGAGGAGACCAGGAGG-3'	21
OPR 215 5'-CCC CCC ACC GGT TAC GAC AAC CCC GTG GTG-3'	22
OPR 216 CCC CCC ATC GAT AAG TGT ACT TAC GTG GGA GAG GGC TTG GAG CAT-3'	23

FIG. 10A

FIG. 10A

APPROVED	D.R. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

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OVA 323-399	
ISQAVHAAHAEINEAGR	26
Gd-246-261	
APYSTLLPPELSETP	27
MBP (83-102) Y83	
YDENPVVHFFKNIVTPRTPP	28
14 amino acid linker	
TSGGGGSGGGGSSS	29
EE TAG	
EEEEYMPMEPG	30
24 amino acid linker	
TSGGGGSGGGGSGGGGSGGGGSSS	31
MBP (S4-102)	
DENPVVHFFKNIVTPRTPP	32

FIG. 10B